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(54) Title: ECIST MICROARRAYS FOR DUAL SCREENING OF DNA HYPERMETHYLATION AND GENE SILENCING

(57) Abstract: Novel methods are herein provided for high-throughput, dual analysis of DNA methylation and gene expression, and triple analysis of DNA methylation, gene expression and gene-associated histone acetylation in cancer cells using arrayed expressed CpG island sequence tags (ECISTs). ECISTs correspond to genomic DNA fragments comprising GC-rich segments along with promoter and/or exon (e.g., first exon) portions of genes. The GC-rich segments are useful for screening hypermethylated CpG sites in cancer cells, while the corresponding promoter and exon-containing portions are useful for determining corresponding transcript levels and assessing histone acetylation. Also provided are high-throughput methods for either confirming methylation-dependent gene silencing, or identifying therapeutically effective demethylating agents, using the ECIST array panels to identify hypermethylated loci, and measure expression levels thereof after cellular exposure to demethylating agents. Further provided are high-throughput methods for distinguishing between direct (primary) demethylation-dependent gene up-regulation, and indirect (secondary) demethylation-dependent up-regulation within apparent epigenetic cascades.

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US03/11598

A. CLASSIFICATION OF SUBJECT MATTER IPC(7) : C12Q 1/00, 1/68; C12P 19/34; C07H 21/02 21/04 US CL : 435/4, 6, 91.1: 536/23.1, 23.5, 24.1, 24.3, 24.33				
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Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched				
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used) Please See Continuation Sheet				
	UMENTS CONSIDERED TO BE RELEVANT			
Category *	Citation of document, with indication, where a		Relevant to claim No.	
Y	HUANG et al. Methylation Profiling of CpG Islam Hum. Mol. Gen. 1999, Vol. 8, No. 3, pages 459-4	70, see whole document.	1-52	
Y	YAN et al. CpG Island Arrays: An Application T Signatures of Breast Cancer. Clin. Canc. Res. Ap whole document.	oward Deciphering Epigenetic ril 2000, Vol. 6, pages 1432-1438, see	1-52	
Y	AHLUWALIA et al. DNA Methylation and Ovarian Cancer: I. Analysis of CpG Island Hypermethylation in Human Ovarian Cancer Using Differential Methylation			
	Hybridazatiion. Gynecologic Oncology. 2001, Vo document.	1. 82, No. 2, pages 261-268, see whole		
YAN et al. Dissecting Complex Epigenetic Alterations in Breast Cancer Using CpG Island Microarays. Cancer Res. 01 December 2001, Vol. 61, pages 8375-8380, see whole document.		1-52		
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	documents are listed in the continuation of Box C.	See patent family annex.		
"A" document	pecial categories of cited documents: defining the general state of the art which is not considered to be lar relevance	"T" later document published after the inter date and not in conflict with the applica principle or theory underlying the inver-	ation but cited to understand the	
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27 July 2003 (27.07.2003) Name and mailing address of the ISA/US (Authorized officer)				
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INTERNATIONAL SEARCH REPORT

PCT/US03/11598

Continuation of Item 4 of the first sheet: The title is too long, PCT Rule 4.3, suggested new title follows: "ECIST MICROARRAYS FOR DUAL SCREENING OF DNA HYPERMETHYLATION AND GENE SILENCING".				
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Continuation of B. FIELDS SEARCHED Item 3: Databases: USPAT, PGPUB, JPO, EPO, Derwent, Medline, Biosis, F	imbara CaDhua Biotachda			
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